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File

 1: AA432248. zw78b02.s1 Soares...[gi:2114636]

Links

IDENTIFIERS

dbEST Id: 1075499
EST name: zw78b02.s1
GenBank Acc: AA432248
GenBank gi: 2114636

CLONE INFO

Clone Id: IMAGE:782283 (3')
Source: IMAGE Consortium, LLNL
DNA type: cDNA

PRIMERS

Sequencing: -41m13 fwd. ET from Amersham
PolyA Tail: Unknown

SEQUENCE

```
TTTTTTTTTTTTCAAAAATGCTTAATAAGTGGACAAACACTGTTGCAA
AATGTAAAGATACTATAAAATTCTTAATACAAAAAGATAAAATTAAAAGCAGATTCTT
TTTTAATTCTGCAACTTGCTCACACGTACATCTTCATTGATTACAGTTGAACAG
AATCCAGTAAATCATTACATGCTCTACAGTCAGTTCAAGGAGCAACCTAACATT
TCCCCCATTATTAAACTAGAGTCATTACACAACCTGTAATAAACTATTGACATTAAT
GTATATGTAACACCTAGTTAACTAAGCAGTAACGGTCATCTGATAGCACCTG
GATGGGGTTGCTATATTAGAACTAAACTAATGAAATGAAAACAAATTGGAATT
A
```

Entry Created: May 22 1997
Last Updated: May 22 1997

COMMENTS

This clone is available royalty-free through LLNL ; contact the IMAGE Consortium (info@image.llnl.gov) for further information.

LIBRARY

Lib Name: Soares_testis_NHT
Organism: Homo sapiens
Sex: male
Lab host: DH10B
Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker
R. Site 1: Not I
R. Site 2: Eco RI
Description: 1st strand cDNA was prepared from mRNA obtained from Clontech Laboratories, Inc., and primed with a Not I - oligo(dT) primer [5' TGTACCAATCTGAAGTGGAGCGGCCGCCAATTTTTTTTTTT 3']. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I

and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization to Cot5, and was constructed by Bento Soares and M. Fatima Bonaldo.

SUBMITTER

Name: Wilson RK
Institution: Washington University School of Medicine
Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
Tel: 314 286 1800
Fax: 314 286 1810
E-mail: est@watson.wustl.edu

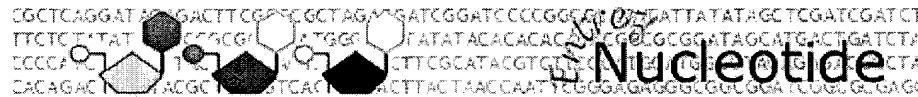
CITATIONS

Title: WashU-Merck EST Project 1997
Authors: Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost ,S., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin ,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising ,B., White,Y., Wylie,T., Waterston,R., Wilson,R.
Year: 1997
Status: Unpublished

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Jun 19 2003 12:37:45



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<input type="button" value="Limits"/>		<input type="button" value="Preview/Index"/>		<input type="button" value="History"/>		<input type="button" value="Clipboard"/>		<input type="button" value="Details"/>
<input type="button" value="Display"/> <input type="button" value="default"/> <input type="button" value="▼"/>		Show: <input type="text" value="20"/> <input type="button" value="▼"/>		<input type="button" value="Send to"/> <input type="button" value="File"/> <input type="button" value="▼"/>				

1: H18957. ym46g08.s1 Soares...[gi:885197]

Links

IDENTIFIERS

dbEST Id: 278895
EST name: ym46g08.s1
GenBank Acc: H18957
GenBank gi: 885197
GDB Id: 424323

CLONE INFO

Clone Id: IMAGE:51513 (3')
DNA type: cDNA

PRIMERS

Sequencing: Promega -21m13
PolyA Tail: Unknown

SEQUENCE

TTTTTTTTGTAAACACTTATTATAAAAAAGTACATTTAATCCTCAGTACATT
 TTCAACCCATCATTTTTTAATACAAGTAAAAGGGGGTATGCAAACACCCCCCAGGT
 CAGAACCGAGGAGATCTGCTGGCTGCCCTGGACCAAGGCAGGAAGGGCGACAAGACG
 CCGAAGCAAGGTAGCGCATCACGCTGGGAGGGAGGGCTGGCAGCTCTCCTGGATTCTT
 TTCATTATACAAAAAAGGAAAACCAATTTTCGACCAAGAATCCCATTCCACAGCA
 GGGTCAGAAGAGCAGCAGCACCGAGT

Quality: High quality sequence stops at base: 305

Entry Created: Jun 29 1995
Last Updated: Jun 29 1995

COMMENTS

High quality sequence stops: 305
 Source: IMAGE Consortium, LLNL
 This clone is available royalty-free through LLNL ; contact
 the IMAGE Consortium (info@image.llnl.gov) for further
 information.

LIBRARY

Lib Name: Soares infant brain 1NIB
Organism: Homo sapiens
Sex: female
Organ: whole brain
Develop. stage: 73 days post natal
Lab host: DH10B (ampicillin resistant)
Vector: Lafmid BA
R. Site 1: Not I
R. Site 2: Hind III
Description: 1st strand cDNA was primed with a Not I - oligo(dT) primer
 [5' AACTGGAAGAATT CGCGGCCAGGAATTTTTTTTTTT 3'];
 double-stranded cDNA was ligated to Hind III adaptors

(Pharmacia), digested with Not I and directionally cloned into the Not I and Hind III sites of the Lafmid BA vector. Library went through one round of normalization. Library constructed by Bento Soares and M.Fatima Bonaldo.

SUBMITTER

Name: Wilson RK
Institution: Washington University School of Medicine
Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
Tel: 314 286 1800
Fax: 314 286 1810
E-mail: est@watson.wustl.edu

CITATIONS

Title: The WashU-Merck EST Project
Authors: Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M.,
Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M.,
, Parsons,J., Rifkin,L., Rohlffing,T., Soares,M., Tan,F.,
Treviskis,E., Waterston,R., Williamson,A., Wohldmann,P.,
Wilson,R.
Year: 1995
Status: Unpublished

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Quackenbush, J.

Year: 2000

Status: Unpublished

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Description: 1st strand cDNA was primed with a Pac I - oligo(dT) primer [5' AACTGGAAAGAATTAAATTAAAGATCTTTTTTTTTTTTT 3'], double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Pac I and cloned into the Pac I and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization. Library constructed by Bento Soares and M.Fatima Bonaldo.

SUBMITTER

Name: Wilson RK
Institution: Washington University School of Medicine
Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
Tel: 314 286 1800
Fax: 314 286 1810
E-mail: est@watson.wustl.edu

CITATIONS

Title: The WashU-Merck EST Project
Authors: Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J., Rifkin,L., Rohlffing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P., Wilson,R.
Year: 1995
Status: Unpublished

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Limits			Preview/Index		History		Clipboard	Details
Display <input type="text" value="default"/>		Show: <input type="text" value="20"/>	Send to <input type="text"/>	File <input type="text"/>				

□1: AA410986. zv03a05.s1 Soares...[gi:2070092]

Links

IDENTIFIERS

dbEST Id: 1055547
EST name: zv03a05.s1
GenBank Acc: AA410986
GenBank gi: 2070092
GDB Id: 5975360

CLONE INFO

Clone Id: IMAGE:752528 (3')
Source: IMAGE Consortium, LLNL
DNA type: cDNA

PRIMERS

Sequencing: -41m13 fwd. ET from Amersham
PolyA Tail: Unknown

SEQUENCE

TTTTTTTTTTTTTGAGGGTAGAGCAGCATCTATTAATATAATTTTATATAGAA
ATACAGGCATATTAAAAATGGAAACATGTAAGAAAGTATGTCACAAGGAATAACAAAAT
ATATCACAAAATAAAAAGTAACCCCAAGTAACAAGTTACTAACACAAGACCCAGCACC
ATGTTGGACTTCTTGATAAGTTCCAGGATGCCAGGCACCTACCGAGGGAGAGATGATC
CTGCTTTGGGAGAGCCAGATGGTCGTGCAGTGGTAAAACCCAGTCCTCCTTTCTGG
AACAGTTATGTCCTGCAAGAAGAAAGACATTGACATCGGGTTGGGCTTACACATTTC
CACCATTTAACAGGGAGACTGGCAAACAGCTGTATGACAGCGAGACCCGTCCAGCCAG
GCGGGCAGTCACACTG

Quality: High quality sequence stops at base: 185

Entry Created: May 2 1997
Last Updated: May 18 1997

COMMENTS

This clone is available royalty-free through LLNL ; contact the IMAGE Consortium (info@image.llnl.gov) for further information.

Possible reversed clone: similarity on wrong strand

PUTATIVE ID Assigned by submitter
TR:G642072 G642072 FIBRILLIN-1

LIBRARY

Lib Name: Soares_NhHMPu_S1
Organism: Homo sapiens
Organ: mixed (see below)
Tissue type: Pooled human melanocyte, fetal heart, and pregnant uterus
Lab host: DH10B
Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker
R. Site 1: Not I

R. Site 2: Eco RI
Description: Equal amounts of plasmid DNA from three normalized libraries (melanocyte 2NbHM, pregnant uterus NbHPU, and fetal heart NbHH19W) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 260232-265223, 340488-345479, and 484488-489479.

SUBMITTER

Name: Wilson RK
Institution: Washington University School of Medicine
Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
Tel: 314 286 1800
Fax: 314 286 1810
E-mail: est@watson.wustl.edu

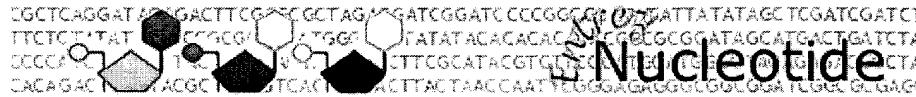
CITATIONS

Title: WashU-Merck EST Project 1997
Authors: Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost ,S., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin ,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising ,B., White,Y., Wylie,T., Waterston,R., Wilson,R.
Year: 1997
Status: Unpublished

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Limits			Preview/Index		History		Clipboard	Details
Display	default	Show: 20	Send to	File				

1: AA620885. af95g06.s1 Soares...[gi:2524824]

Links

IDENTIFIERS

dbEST Id: **1318692**
 EST name: af95g06.s1
 GenBank Acc: AA620885
 GenBank gi: 2524824

CLONE INFO

Clone Id: IMAGE:1055578 (3')
 Source: IMAGE Consortium, LLNL
 Insert length: 1052
 DNA type: cDNA

PRIMERS

Sequencing: -40m13 fwd. ET from Amersham
 PolyA Tail: Unknown

SEQUENCE

TTTGAACATAAAAATTCTTATTTAACCTAACGAGCAGTATTGAGATAGTTGCTATA
 TTAAAAACAAGACGTTAAAAAAATTACAGCAAAGTTAGCAAGGCAGTGACTAATTAAGT
 CACTAAGTTAATTATATTCTTCACAGTCATTCTATAATCATGTAATGGTAAACAATA
 TTTTCAGCCACTTGGAGATAAGTTAACATTAAAGTTCCAGAAATAACACAGTAAAGAAA
 TTGAATTATATAAAAGAGGTTAAACATTAAAGTTCCAGAAATAACACAGTAAAGAAA
 TATGAAAATAAACTGGAAAATAAAATACCCACCCATCCGAAAATCTACATCATCTCT
 TTCATTGTCCCCAATGCCTTTC

Quality: High quality sequence stops at base: 344

Entry Created: Oct 15 1997
 Last Updated: Mar 2 1998

COMMENTS

This clone is available royalty-free through LLNL ; contact the IMAGE Consortium (info@image.llnl.gov) for further information.

LIBRARY

Lib Name: Soares_testis_NHT
 Organism: Homo sapiens
 Sex: male
 Lab host: DH10B
 Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker
 R. Site 1: Not I
 R. Site 2: Eco RI
 Description: 1st strand cDNA was prepared from mRNA obtained from Clontech Laboratories, Inc., and primed with a Not I - oligo(dT) primer [5' TGTTACCAATCTGAAGTGGGAGCGGCCGCCAATTTTTTTTTTT 3']. Double-stranded cDNA was ligated to Eco RI adaptors

(Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization to Cot5, and was constructed by Bento Soares and M. Fatima Bonaldo.

SUBMITTER

Name: Wilson RK
Institution: Washington University School of Medicine
Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
Tel: 314 286 1800
Fax: 314 286 1810
E-mail: est@watson.wustl.edu

CITATIONS

Title: WashU-NCI human EST Project
Authors: Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost ,S., Krizman,D., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra ,M., Martin,J., Moore,B., Schellenberg,K., Steptoe,M., Tan ,F., Theising,B., White,Y., Wylie,T., Waterston,R., Wilson ,R.
Year: 1997
Status: Unpublished

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Limits		Preview/Index		History		Clipboard	Details	
Display <input type="button" value="default"/> <input type="button" value="▼"/>		Show: <input type="button" value="20"/> <input type="button" value="▼"/>		<input type="button" value="Send to"/> <input type="button" value="File"/> <input type="button" value="▼"/>				

□ 1: AI368878. qw15a11.x1 NCI CG...[gi:4147631]

Links

IDENTIFIERS

dbEST Id: 2157828
EST name: qw15a11.x1
GenBank Acc: AI368878
GenBank qi: 4147631

CLONE INFO

Clone Id: IMAGE:1991132 (3')
Source: NCI
Insert length: 3608
DNA type: cDNA

PRIMERS

Sequencing: -40UP from Gibco
PolyA Tail: Unknown

SEQUENCE

TTTTTTTAAACATAAAAATTCTTTATTAACCTAATCCAG
TATATTAAAAACAAGACGTTAAAAAAATTACAGCAAAGT
AAGTCACTAAGTTAATTATATTCTTCACAGTCATTC
AATATTTCAGGCCACTTGAGAGATAAGTTAACCTTGAAA
GTCATTGAATTTATAAAAGAGGTTAAACATTAAGTT
GAAATATGAAAATAAACTGGAAAATAAAATATACCCACCC
CTCTTCATTTGTCCCCATGCCTTCTTTACCTCTCCA
Quality: High quality sequence stops at base: 399

Entry Created: Jan 11 1999
Last Updated: Feb 15 1999

COMMENTS

Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,
Michael R. Emmert-Buck, M.D., Ph.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing
Center
Clone distribution: NCI-CGAP clone distribution information
can be found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html

LIBRARY

Lib Name: NCI_CGAP_Ut3
Organism: Homo sapiens
Organ: uterus
Tissue type: poorly-differentiated endometrial adenocarcinoma, 2 pooled tumors
Lab host: DH10B

Vector: pCMV-SPORT6
R. Site 1: Sall
R. Site 2: NotI
Description: Cloned unidirectionally. Primer: Oligo dT. Average insert size 1.45 kb. Life Technologies catalog #: 11541-018

SUBMITTER

Name: Robert Strausberg, Ph.D.
E-mail: cgapbs-r@mail.nih.gov

CITATIONS

Title: National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Tumor Gene Index
Authors: NCI-CGAP <http://www.ncbi.nlm.nih.gov/ncicgap>
Year: 1997
Status: Unpublished

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Description: 1st strand cDNA was primed with a Not I - oligo(dT) primer [5' TGTTACCAATCTGAAGTGGGAGCGGCCGCATTTTTTTTTTTTT 3'], double-stranded cDNA was size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector (Pharmacia). Library went through one round of normalization to a Cot = 5. Library constructed by Bento Soares and M.Fatima Bonaldo. RNA from 4 multiple sclerosis lesions from one patient was kindly provided by Dr. Kevin G. Becker (NINDS/NIH).

SUBMITTER

Name: Wilson RK
Institution: Washington University School of Medicine
Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
Tel: 314 286 1800
Fax: 314 286 1810
E-mail: est@watson.wustl.edu

CITATIONS

Title: The WashU-Merck EST Project
Authors: Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M.,
Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M.,
Parsons,J., Rifkin,L., Rohlffing,T., Soares,M., Tan,F.,
Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P.,
Wilson,R.
Year: 1995
Status: Unpublished

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GGCTCAGGATAGGACTTCC>CGCTAG>AGATCGGATCCCCGGCG>TGATTATATAGCTCGATCGATCT
TTCTCT>TAT>CGCGG>ATGC>ATATACACACAC>CGGGGGATAGCATGACT>GATCT>
CCCC>A>T>G>C>TTCGCGATACGTGCG>CGATGATG>ATG>CGATGATCT
GAGAGAC>AGC>TCAT>ACTTAACTAAC>TGGGAGGGG>GGGGATGAGGAG

PubMed Nucleotide Protein Genome Structure PMC Taxonomy OMIM Books

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Show: 20

1: H98215.yx09a02.s1 Soares...[gi:1119100]

Links

IDENTIFIERS

dbEST Id: 407094
EST name: yx09a02.s1
GenBank Acc: H98215
GenBank gi: 1119100
GDB Id: 3870836

CLONE INFO

Clone Id: IMAGE:261194 (3')
Insert length: 1053
DNA type: cDNA

PRIMERS

Sequencing: m13 -40 forward
PolyA Tail: Unknown

SEQUENCE

CACAGGAACAATTCTTTATTGTACATTGGAGAAATAGCCCTGTGTGCTGGTTCAAGGTG
 CAACATACAGAATATTGAATTAAGAAAAGAGGGAACGGGAAGGGAAANGGAAACCTCTT
 GAGGTCAAAGTTGNCACAAAAATGGTAAAGAAAAGATTTCCTCACGCAAGANGGCATTTT
 GCAAATACCATGCAAAACAGGCAGCTGGTGTGCCTTAAGAGAATCCCTATAATAACAGA
 AAAGACACTCCAAGCATTCTGTACGTGGACTCAGAGCACAGAGAAAAGAAACTAAATG
 CCTTTGGATTCAAGATATTGGACTCTTGATTACATTTTTACAGTCCATTAAA
 GGGGAATAAACTGACATAATATT

Quality: High quality sequence stops at base: 109

Entry Created: Dec 12 1995
Last Updated: Dec 12 1995

COMMENTS

High quality sequence stops: 109
 Source: IMAGE Consortium, LLNL
 This clone is available royalty-free through LLNL ; contact
 the IMAGE Consortium (info@image.llnl.gov) for further
 information.

LIBRARY

Lib Name: Soares melanocyte 2NbHM
Organism: Homo sapiens
Sex: Male
Tissue type: melanocyte
Lab host: DH10B (ampicillin resistant)
Vector: pT7T3D (Pharmacia) with a modified polylinker
R. Site 1: Not I
R. Site 2: Eco RI
Description: 1st strand cDNA was primed with a Not I - oligo(dT) primer
 [5' TGTTACCAATCTGAAGTGGAGCGGCCGAGTTTTTTTTTT 3'],

double-stranded cDNA was size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector (Pharmacia). Library constructed by Bento Soares and M.Fatima Bonaldo. RNA from normal foreskin melanocytes (FS374) was kindly provided by Dr. Anthony P. Albino.

SUBMITTER

Name: Wilson RK
Institution: Washington University School of Medicine
Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
Tel: 314 286 1800
Fax: 314 286 1810
E-mail: est@watson.wustl.edu

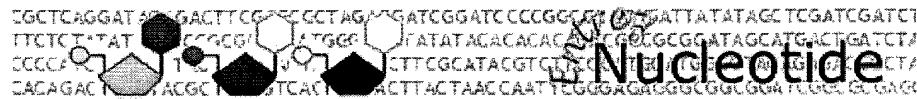
CITATIONS

Title: The WashU-Merck EST Project
Authors: Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M.,
Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M.,
, Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F.,
Trevisakis,E., Waterston,R., Williamson,A., Wohldmann,P.,
Wilson,R.
Year: 1995
Status: Unpublished

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Search <input type="text" value="Nucleotide"/> <input checked="" type="checkbox"/> for <input type="text"/>				<input type="button" value="Go"/> <input type="button" value="Clear"/>				
Limits		Preview/Index			History		Clipboard	Details
<input type="button" value="Display"/> <input type="button" value="default"/>		Show: <input type="text" value="20"/> <input type="button"/>		<input type="button" value="Send to"/> <input type="button" value="File"/>				

1: T87560. yd83b10.r1 Soares...[gi:715912]

Links

IDENTIFIERS

dbEST Id: 157681
EST name: yd83b10.r1
GenBank Acc: T87560
GenBank gi: 715912
GDB Id: 470428

CLONE INFO

Clone Id: IMAGE:114811 (5')
Insert length: 1110
DNA type: cDNA

PRIMERS

Sequencing: M13RP1
PolyA Tail: Unknown

SEQUENCE

NTAATACTGAGAGCGGAAGATTTTAAAGAGATATGGTAGTTAAATAGGTTGTTA
 AAGAACTCTGGAAATTCTAACAAATTGTAATTGTTTATAAACCTCCTACAGAG
 TTGTGTACCATTCATCATCACCATCATTAAATATCTGTAAGTGCCCAGAAGGTACAAGGT
 GCTATACAGCCGTTAACACCCCCCTCCCTGCAACTGTTAAAATTAGAATAGATTTTGC
 ATCAACCCTAGCTAGAAGCTTCCCCACCTAGGTGGCCTGGCAAGGGCCTGCATTCCCAA
 CATCCCATACATGTTCCCACCTCCAGAAAGTCTCTGGGTTATTCCAGCNTTTTATC
 NTGGACGTGTCCTTCACCCCTGGGTTGGGTTGGGTTGTTAAANTC
 AGGAAAGTTTCAGGGATTCAAAGGGGTCNGGACNTTTTAACCCGGCGAGTACAGG
 TCCNTTTTGCTANGTTGTTGTGGTACAGGNAACCCANTGGCNTTAGTGTGT
 GGGTAGTTGT

Quality: High quality sequence stops at base: 324

Entry Created: Mar 17 1995
Last Updated: Mar 17 1995

COMMENTS

Insert Size: 1110
 High quality sequence stops: 324 Source: IMAGE Consortium,
 LLNL This clone is available royalty-free through LLNL ;
 contact the IMAGE Consortium (info@image.llnl.gov) for
 further information.

LIBRARY

Lib Name: Soares fetal liver spleen 1NFLS
Organism: Homo sapiens
Sex: male
Organ: Liver and Spleen
Develop. stage: 20 week-post conception fetus
Lab host: DH10B (ampicillin resistant)
Vector: pT7T3D (Pharmacia) with a modified polylinker

R. Site 1: Pac I
R. Site 2: Eco RI
Description: 1st strand cDNA was primed with a Pac I - oligo(dT) primer [5' AACTGGAAGAATTAAATTAAAGATCTTTTTTTTTTTTT 3'], double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Pac I and cloned into the Pac I and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization. Library constructed by Bento Soares and M.Fatima Bonaldo.

SUBMITTER

Name: Wilson RK
Institution: Washington University School of Medicine
Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
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Fax: 314 286 1810
E-mail: est@watson.wustl.edu

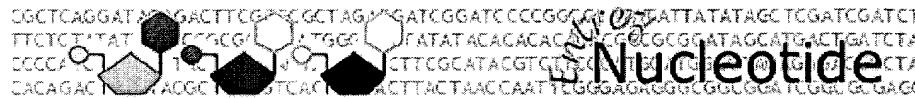
CITATIONS

Title: The WashU-Merck EST Project
Authors: Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J., Rifkin,L., Rohlffing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P., Wilson,R.
Year: 1995
Status: Unpublished

MAP DATA

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 1: AW275150. xv76c12.x1 NCI_CG...[gi:6662180]

Links

IDENTIFIERS

dbEST Id: 3658251
EST name: xv76c12.x1
GenBank Acc: AW275150
GenBank gi: 6662180

CLONE INFO

Clone Id: IMAGE:2819062 (3')
Source: NCI
DNA type: cDNA

PRIMERS

Sequencing: -40UP from Gibco
PolyA Tail: Unknown

SEQUENCE

TTTTTTGAGGAATGAGCAATTATTCAATAAAAGAGAAAGCATTAACTTGTACAGTG
GGAAAAAAATGAACCTCAAGAGTTGCTACATTAACTGTATCCCCATTATCTGCACGAT
GTCTTATCTCAGTGTCTAACACTAAAATATTGAATGAGAAATACACCAACGTTGG
CTGATTGCTTGACATGTCGATTAGGGAGACTTCTACAACCACTCCTCTTTCTC
CCAGTAAATACTTTGACTTGACACCTACCATATTGAAATGACAGGTGCCGAGGGCA
AGTGCATCAAAGCAGTTAGGATTGCAATGCTGTAAGGATTATTTTTAATGGAGCAG
TTCTATTGAATCCTGAAATGCAGAACCCAGTGAATCTCCTTAT

Quality: High quality sequence stops at base: 404

Entry Created: Jan 3 2000
Last Updated: Jan 3 2000

COMMENTS

Tissue Procurement: Chris Moskaluk, M.D., Ph.D., Michael R. Emmert-Buck, M.D., Ph.D. cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: Christa Prange, The I.M.A.G.E. Consortium DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html

LIBRARY

Lib Name: NCI_CGAP_Lu28
Organism: Homo sapiens
Organ: lung
Tissue type: two pooled squamous cell carcinomas
Lab host: DH10B
Vector: pCMV-SPORT6
R. Site 1: SalI
R. Site 2: NotI

Description: Cloned unidirectionally. Primer: Oligo dT. Library constructed by Life Technologies.

SUBMITTER

Name: Robert Strausberg, Ph.D.
E-mail: cgapbs-r@mail.nih.gov

CITATIONS

Title: National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Tumor Gene Index
Authors: NCI-CGAP <http://www.ncbi.nlm.nih.gov/ncicgap>
Year: 1997
Status: Unpublished

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The diagram illustrates a segment of a nucleotide strand. The backbone consists of alternating phosphate groups (represented by orange and red hexagons) and deoxyribose sugar rings (represented by grey pentagons). Four nitrogenous bases are attached to the sugars: Adenine (A), Thymine (T), Cytosine (C), and Guanine (G). Each base has specific chemical substituents: Adenine has an amino group (-NH₂) and a methyl group (-CH₃); Thymine has a methyl group (-CH₃); Cytosine has a carbonyl group (-C=O) and a methyl group (-CH₃); and Guanine has two amino groups (-NH₂). The entire structure is labeled "Nucleotide".

PubMed	Nucleotide	Protein	Genome	Structure	PMC	Taxonomy	OMIM	Books
Search <input type="text" value="Nucleotide"/> for <input type="text"/>						<input type="button" value="Go"/>	<input type="button" value="Clear"/>	
Limits		Preview/Index		History		Clipboard	Details	
Display <input type="text" value="default"/> <input type="button" value="▼"/>		Show: <input type="text" value="20"/> <input type="button" value="▼"/>	Send to <input type="text" value="File"/> <input type="button" value="▼"/>					

□ 1: AI521564. to64g08.x1 NCI CG...[gi:4435699]

Links

IDENTIFIERS

dbEST Id: 2337724
EST name: to64g08.x1
GenBank Acc: AI521564
GenBank gi: 4435699

CLONE INFO

Clone Id: IMAGE:2183102 (3')
Source: NCI
Insert length: 4578
DNA type: cDNA

PRIMERS

Sequencing: -40UP from Gibco
PolyA Tail: no

SEQUENCE

Quality: High quality sequence stops at base: 371

AGCGGCCGCCCCTTTTTTTTTTTGTGAGGAATGAGCAATTATTCATAAAG
AGAAAAGCATTAAATTTCGTACAGTGGGAAAAAATGAACCTAACAGAGTGCTACATTAACT
GTATCCCATTATCTCTGCACGATGTCTTATCTCAGTGTCTCAATTCACACTAAAATA
TTGAATGAGAAATAACACCACTGGCTGATTGCTGACATGCTGATTAGGGAGACTTC
TACAACCACCTCTCTTTCTCCAGTAAAATACTTTGACTTTGACACCTACCATAT
TGGAAATGACAGGTGCCCGAGGGCAAGTGCATCAAAGCAGTTAGGATTCGAATGCTTGCT
AAGGATTATTTTTAATGGAGCAGTTCTATTGAATCTGAAAATGCAGAACCCAG

Quality: High quality sequence stops at base: 371

Entry Created: Mar 16 1999
Last Updated: May 12 1999

COMMENTS

Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,
Michael R. Emmert-Buck, M.D., Ph.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing
Center
Clone distribution: NCI-CGAP clone distribution information
can be found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html

LIBRARY

Lib Name: NCI_CGAP_Gas4
Organism: Homo sapiens
Organ: stomach
Tissue type: poorly differentiated adenocarcinoma with signet ring cell features
Lab host: DH10B

Vector: pCMV-SPORT6
R. Site 1: SalI
R. Site 2: NotI
Description: Cloned unidirectionally. Primer: Oligo dT. Average insert size 1.69 kb. Life Technologies catalog #: 11549-011

SUBMITTER

Name: Robert Strausberg, Ph.D.
E-mail: cgapbs-r@mail.nih.gov

CITATIONS

Title: National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Tumor Gene Index
Authors: NCI-CGAP <http://www.ncbi.nlm.nih.gov/ncicgap>
Year: 1997
Status: Unpublished

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